

REPLACEMENT SHEET

Fig. 6D

ACAGCGACTCAGATTTCAGACAGCGACTCGGATTCTGATAGCGACTCGGATTTCAGACAGCG-2640
D S D S D S D S D S D S D S D S D S D S -869

ACTCAGACTCAGACAGTGATTTCAGATTTCAGACAGCGACTCAGATTCCGATAGTGATTTCAG-2700
D S D S D S D S D S D S D S D S D S D S -889

ACTCAGACAGCGACTCAGATTCTGATAGTGATTTCAGACTCAGACAGTGATTTCAGATTTCAG-2760
D S D S D S D S D S D S D S D S D S D S -909

ACAGCGACTCAGATTCCGATAGTGATTTCAGACTCAGACAGCGACTCAGATTCCGATAGTG-2820
D S D S D S D S D S D S D S D S D S D S -929

ATTCAGACTCAGACAGCGACTCAGATTCTGATAGTGATTTCAGACTCAGACAGTGATTTCAG-2880
D S D S D S D S D S D S D S D S D S D S -949

ACTCAGACAGTGATTTCAGATTCCGATAGTGATTTCAGACTCCGATAGCGACTCAGACTCGG-2940
D S D S D S D S D S D S D S D S D S D S -969

ATAGTGACTCAGATTCTGATAGTGATTTCAGACTCCGATAGCGACTCAGACTCCGATAGTG-3000
D S D S D S D S D S D S D S D S D S D S -989

ACTCAGATTCTGATAGTGATTTCAGACTCAGACAGCGACTCAGATTCTGATAGTGATTTCAG-3060
D S D S D S D S D S D S D S D S D S D S -1009

ACTCAGTCAGTGATTTCAGATTCCGATAGTGATTTCAGACTCAGGCAGTGATTCCGATTCCG-3120
D S V S D S D S D S D S D S G S D S D S -1029

R←

ATAGTGATTTCAGACTCAGACAACGACTCAGATTTAGGCAATAGCTCAGATAAGAGTACAA-3180
D S D S D S D N D S D L G N S S D K S T -1049

→M

AAGATAAATTACCTGATACAGGAGCTAATGAAGATTATGGCTCTAAAGGCACGTTACTTG-3240
K D K L P D T G A N E D Y G S K G T L L -1069

GAACTCTGTTTGCAGGTTTGGAGCGTTATTATTAGGGAAACGTCGCAAAAATAGAAAA-3300
G T L F A G L G A L L L G K R R K N R K -1089

ATAAAAATTAAATGTTCAAATGAAATTTGTAGAAAGAAGCAGATATGAGATTTGAATAG-3360
N K N * -1092

AAAGTAGATTTAGTCCAACAAATGTAAGATGTTGATTAAACTATAATATAACTTTTCAG-3420

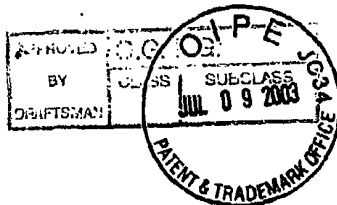


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ACAGCGACTCAGATTCAGACAGCGACTCGGATTCTGATAGCGACTCGGATTCAGACAGCG-2640
D S D S D S D S D S D S D S D S D S -869

ACTCAGACTCAGACAGTGATTCAGATTCAGACAGCGACTCAGATTCGATAGTGATTCAG-2700
D S D S D S D S D S D S D S D S D S -889

ACTCAGACAGCGACTCAGATTCGATAGTGATTCAGACTCAGACAGTGATTCAGATTCAG-2760
D S D S D S D S D S D S D S D S D S -909

ACAGCGACTCAGATTCGATAGTGATTCAGACTCAGACAGCGACTCAGATTCGATAGTG-2820
D S D S D S D S D S D S D S D S D S -929

ATTCAGACTCAGACAGCGACTCAGATTCGATAGTGATTCAGACTCAGACAGTGATTCAG-2880
D S D S D S D S D S D S D S D S D S -949

ACTCAGACAGTGATTCAGATTCGATAGTGATTCAGACTCCGATAGCGACTCAGACTCGG-2940
D S D S D S D S D S D S D S D S D S -969

ATAGTGAATCAGATTCGATAGTGATTCAGACTCCGATAGCGACTCAGACTCGGATAGTG-3000
D S D S D S D S D S D S D S D S D S -989

ACTCAGATTCGATAGTGATTCAGACTCAGACAGCGACTCAGATTCGATAGTGATTCAG-3060
D S D S D S D S D S D S D S D S D S -1009

ACTCAGTCAGTGATTCAGATTCGATAGTGATTCAGACTCAGGCAGTGATTCGGATTCGG-3120
D S V S D S D S D S D S D S G S D S D S -1029

R←
ATAGTGATTCAGACTCAGACAACGACTCAGATTTAGGCAATAGCTCAGATAAGAGTACAA-3180
D S D S D S D N D S D L G N S S D K S T -1049

→M
AAGATAAATTACCTGATAGAGGAGCTAATGAAGATTATGGCTCTAAAGGCACGTTACTTG-3240
K D K L P D T G A N E D Y G S K G T L L -1069

GAACTCTGTTTGCGGTTTAGGAGCGTTATTATTAGGGAAACGTCGCAAAAATAGAAAAA-3300
G T L F A G L G A L L L G K R R K N R K -1089

ATAAAAATTAAATGTTCAAATGAAATTTGTAGAAAGAAGCAGATATGAGATTTGAATAG-3360
N K N * -1092

AAAGTAGATTTAGTCCAACAAATGTAAGATGTTGATTAAAACTATAATATAACTTTCACG-3420